RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/540,846
Source: IFW P
Date Processed by STIC: 06/21/2006

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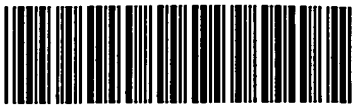

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RAW SEQUENCE LISTING DATE: 06/21/2006
PATENT APPLICATION: **US/10/540,846** TIME: 10:16:53

Input Set : **E:\ARS-111 seq-listing.replace.txt**
Output Set: **N:\CRF4\06212006\J540846.raw**

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 3 <110> APPLICANT: Bienkowska, Jadwiga
 4       Mcallister, Gregg
 6 <120> TITLE OF INVENTION: NOVEL FIBULIN-LIKE POLYPEPTIDES
 8 <130> FILE REFERENCE: ARS-111
10 <140> CURRENT APPLICATION NUMBER: US 10/540,846
11 <141> CURRENT FILING DATE: 2005-06-27
13 <150> PRIOR APPLICATION NUMBER: US 60/436,786
14 <151> PRIOR FILING DATE: 2002-12-27
16 <160> NUMBER OF SEQ ID NOS: 4
18 <170> SOFTWARE: PatentIn version 3.1
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22 <212> TYPE: DNA
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26 <221> NAME/KEY: misc_feature
27 <222> LOCATION: (50)..(2582)
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35 caccgggagg aagccgcccg ggcacttcgc ggccgagaga cgccgactgg gcccccacgt    180
37 ctgcctctct gggtttggga gtggctgctg ccctggctgg gcgccctcta tgggtggtgg    240
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55 tcggaggcca ttggagaggc gagtctgtca ccattcctgc cacaacaccg tgggcagctt    780
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 85 cgtgtgccga gactgcaact acgagggaag gaaggtggcg aatggccagg tgttcacctt   1680
 87 ggatgatgaa ccctgcaccc ggtgcacgtg ccagctggga gaggtgagct gtgagaaggt   1740
 89 tccctgccag cgggcctgtg ccgaccctgc cctgcttcct ggggactgct gctcttcctg   1800
 91 tccagattcc ctgtctcctc tggaagaaaa gcaggggctc tcccctcacg gaaatgtggc   1860
 93 attcagcaaa gctggtcgga gcctgcatgg agacactgag gcccctgtca actgtagctc   1920
 95 ctgtcctggg cccccgacag catcaccctc gaggccggtg cttcatctcc tccagctcct   1980
 97 tttaagaacg aacttgatga aaacacagac tttacctaca agcccggcag gagctcatgg   2040
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103 aggggctcct cagccacctc ctgtgactcc agagcgctcg ttctcagcct ctggggccca   2220
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113 tgcactgctc attcaccgct tacctgtggg aaggtgggaa acgtgacccc aagcccacag   2520
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133             20                  25                  30
135 Phe Ala Ala Glu Arg Arg Arg Leu Gly Pro His Val Cys Leu Ser Gly
136         35                  40                  45
138 Phe Gly Ser Gly Cys Cys Pro Gly Trp Ala Pro Ser Met Gly Gly Gly
139     50                  55                  60
141 His Cys Thr Leu Pro Leu Cys Ser Phe Gly Cys Gly Ser Gly Ile Cys
142 65                  70                  75                  80
144 Ile Ala Pro Asn Val Cys Ser Cys Gln Asp Gly Glu Gln Gly Ala Thr
145                 85                  90                  95
147 Cys Pro Glu Thr His Gly Pro Cys Gly Glu Tyr Gly Cys Asp Leu Thr
148             100                 105                 110
150 Cys Asn His Gly Gly Cys Gln Glu Val Ala Arg Val Cys Pro Val Gly
151         115                 120                 125
153 Phe Ser Met Thr Glu Thr Ala Val Gly Ile Arg Cys Thr Asp Ile Asp
154     130                 135                 140
156 Glu Cys Val Thr Ser Ser Cys Glu Gly His Cys Val Asn Thr Glu Gly
157 145                 150                 155                 160
159 Gly Phe Val Cys Glu Cys Gly Pro Gly Met Gln Leu Ser Ala Asp Arg
160                 165                 170                 175
162 His Ser Cys Gln Asp Thr Asp Glu Cys Leu Gly Thr Pro Cys Gln Gln
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165 Arg Cys Lys Asn Ser Ile Gly Ser Tyr Lys Cys Ser Cys Arg Thr Gly
166         195                 200                 205
168 Phe His Leu His Gly Asn Arg His Ser Cys Val Asp Val Asn Glu Cys
169     210                 215                 220
171 Arg Arg Pro Leu Glu Arg Arg Val Cys His His Ser Cys His Asn Thr
172 225                 230                 235                 240
174 Val Gly Ser Phe Leu Cys Thr Cys Arg Pro Gly Phe Arg Leu Arg Ala
175                 245                 250                 255
177 Asp Arg Val Ser Cys Glu Gly Ala Leu Ser Pro Pro Asp Trp Gln Gln
178                 260                 265                 270
180 Gly Pro Leu Pro Ala Gly Thr Trp Glu Pro Cys Met Asn Gln Gly Val
181         275                 280                 285
183 Ala Gly Gln Ser Leu Gly Val Pro Ser Ala Gly Ala Arg Leu Glu Thr
184     290                 295                 300
186 Cys Arg Ala Cys Phe Val Ser Val Leu Leu Ala Arg Val Arg Pro Pro
187 305                 310                 315                 320
189 Ile Lys Thr Asp Cys Cys Thr Cys Val Pro Val Arg Cys Tyr Phe His
190                 325                 330                 335
192 Gly Arg Trp Tyr Ala Asp Gly Ala Val Phe Ser Gly Gly Gly Asp Glu
193                 340                 345                 350
195 Cys Thr Thr Cys Val Cys Gln Asn Gly Glu Val Glu Cys Ser Phe Met
196         355                 360                 365
198 Pro Cys Pro Glu Leu Ala Cys Pro Arg Glu Glu Trp Arg Leu Gly Pro
199     370                 375                 380
201 Gly Gln Cys Cys Phe Thr Cys Gln Glu Pro Thr Pro Ser Thr Gly Cys
202 385                 390                 395                 400
204 Ser Leu Asp Asp Asn Gly Val Glu Phe Pro Ile Gly Gln Ile Trp Ser
205                 405                 410                 415
207 Pro Gly Asp Pro Cys Glu Leu Cys Ile Cys Gln Ala Asp Gly Ser Val
208                 420                 425                 430
210 Ser Cys Lys Arg Thr Asp Cys Val Asp Ser Cys Pro His Pro Ile Arg
211         435                 440                 445
213 Ile Pro Gly Gln Cys Cys Pro Asp Cys Ser Ala Ala Gly Ala Gln Arg
214     450                 455                 460
216 Met Leu Ser Leu Ala Gly Cys Thr Tyr Thr Gly Arg Ile Phe Tyr Asn
217 465                 470                 475                 480
219 Asn Glu Thr Phe Pro Ser Val Leu Asp Pro Cys Leu Ser Cys Ile Cys
220                 485                 490                 495
222 Leu Leu Gly Ser Val Ala Cys Ser Pro Val Asp Cys Pro Ile Thr Cys
223                 500                 505                 510
225 Thr Tyr Pro Phe His Pro Asp Gly Glu Cys Cys Pro Val Cys Arg Asp
226         515                 520                 525
228 Cys Asn Tyr Glu Gly Arg Lys Val Ala Asn Gly Gln Val Phe Thr Leu
229     530                 535                 540
231 Asp Asp Glu Pro Cys Thr Arg Cys Thr Cys Gln Leu Gly Glu Val Ser
232 545                 550                 555                 560
234 Cys Glu Lys Val Pro Cys Gln Arg Ala Cys Ala Asp Pro Ala Leu Leu
235                 565                 570                 575
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237 Pro Gly Asp Cys Cys Ser Ser Cys Pro Asp Ser Leu Ser Pro Leu Glu
238             580                 585                 590
240 Glu Lys Gln Gly Leu Ser Pro His Gly Asn Val Ala Phe Ser Lys Ala
241         595                 600                 605
243 Gly Arg Ser Leu His Gly Asp Thr Glu Ala Pro Val Asn Cys Ser Ser
244     610                 615                 620
246 Cys Pro Gly Pro Pro Thr Ala Ser Pro Ser Arg Pro Val Leu His Leu
247 625                 630                 635                 640
249 Leu Gln Leu Leu Leu Arg Thr Asn Leu Met Lys Thr Gln Thr Leu Pro
250                 645                 650                 655
252 Thr Ser Pro Ala Gly Ala His Gly Pro His Ser Leu Ala Leu Gly Leu
253             660                 665                 670
255 Thr Ala Thr Phe Pro Gly Glu Pro Gly Ala Ser Pro Arg Leu Ser Pro
256         675                 680                 685
258 Gly Pro Ser Thr Pro Pro Gly Ala Pro Thr Leu Pro Leu Ala Ser Pro
259     690                 695                 700
261 Gly Ala Pro Gln Pro Pro Pro Val Thr Pro Glu Arg Ser Phe Ser Ala
262 705                 710                 715                 720
264 Ser Gly Ala Gln Ile Val Ser Arg Trp Pro Pro Leu Pro Gly Thr Leu
265                 725                 730                 735
267 Leu Thr Glu Ala Ser Ala Leu Ser Met Met Asp Pro Ser Pro Ser Lys
268             740                 745                 750
270 Thr Pro Ile Thr Leu Leu Gly Pro Arg Val Leu Ser Pro Thr Thr Ser
271         755                 760                 765
273 Arg Leu Ser Thr Ala Leu Ala Ala Thr Thr His Pro Gly Pro Gln Gln
274     770                 775                 780
276 Pro Pro Val Gly Ala Ser Arg Gly Glu Glu Ser Thr Met Leu Ser Arg
277 785                 790                 795                 800
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295 atgggtggtg ggcactgcac cctgcccctc tgctccttcg gctgtgggag tggcatctgc    180
297 atcgctccca atgtctgctc ctgccaggat ggagagcaag gggccacctg cccagaaacc    240
299 catggaccat gtggggagta cggctgtgac cttacctgca accatggagg ctgtcaggag    300
301 gtggcccgag tgtgccccgt gggcttctcg atgacggaga cagctgttgg catcaggtgt    360
303 acagacattg acgaatgtgt aacctcctcc tgcgagggcc actgtgtgaa cacagaaggt    420
305 gggtttgtgt gcgagtgtgg gccgggcatg cagctgtctg ccgaccgcca cagctgccaa    480
307 gacactgacg aatgcctagg gactccctgt cagcagagat gtaaaaacag cattggcagc    540
309 tacaagtgtt cctgtcgaac tggcttccac cttcatggca accggcactc ctgtgtagat    600
311 gtaaacgagt gtcggaggcc attggagagg cgagtctgtc accattcctg ccacaacacc    660
313 gtgggcagct tcctatgcac atgccgacct ggcttcaggc tccgagctga ccgcgtgtcc    720
315 tgtgaagggg ccctgagtcc ccccgactgg cagcagggcc ctctccctgc tggcacctgg    780
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321 ataaagacgg attgctgtac ttgtgttcca gtgagatgct atttccacgg ccggtggtac    960
323 gcagacgggg ctgtgttcag tgggggtggt gacgagtgta ccacctgtgt ttgccagaat   1020
325 ggggaggtgg agtgctcctt catgccctgc cctgagctgg cctgcccccg agaagagtgg   1080
327 cggctgggcc ctgggcagtg ttgcttcacc tgccaggagc ccacaccctc gacaggctgc   1140
329 tctcttgacg acaacggggt tgagtttccg attggacaga tctggtcgcc tggtgacccc   1200
331 tgtgagttat gcatctgcca ggcagatggc tcggtgagct gcaagaggac agactgtgtg   1260
333 gactcctgcc ctcacccgat ccggatccct ggacagtgct gcccagactg ttcagcagct   1320
335 ggtgctcagc gcatgctctc tctggcaggc tgcacctaca caggcagaat cttctataac   1380
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351 aactgtagct cctgtcctgg gcccccgaca gcatcaccct cgaggccggt gcttcatctc   1860
353 ctccagctcc ttttaagaac gaacttgatg aaaacacaga ctttacctac aagcccggca   1920
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386         35                  40                  45
388 Pro Leu Cys Ser Phe Gly Cys Gly Ser Gly Ile Cys Ile Ala Pro Asn
389     50                  55                  60
391 Val Cys Ser Cys Gln Asp Gly Glu Gln Gly Ala Thr Cys Pro Glu Thr
392 65                  70                  75                  80
394 His Gly Pro Cys Gly Glu Tyr Gly Cys Asp Leu Thr Cys Asn His Gly
395                 85                  90                  95
397 Gly Cys Gln Glu Val Ala Arg Val Cys Pro Val Gly Phe Ser Met Thr
398             100                 105                 110
400 Glu Thr Ala Val Gly Ile Arg Cys Thr Asp Ile Asp Glu Cys Val Thr
401         115                 120                 125
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